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SEQUENCE LISTING

<110> Bayer AG

<120> NOVEL EIMERIA GENE AND PROTEIN, AND THEIR USE

<130> LeA 36695 DE

<160> 24

<170> PatentIn version 3.1

<210> 1

<211> 1186

<212> DNA

<213> Eimeria tenella

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<221> CDS

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ggctacaaga acaagtttga ag atg agg act atc cta gcc acc cta gtc ggt		112
Met Arg Thr Ile Leu Ala Thr Leu Val Gly		
1	5	10
ttc aca gcc tgc gca gcc gtt gct gca gac gga gca cct gag tat cct		160
Phe Thr Ala Cys Ala Ala Val Ala Ala Asp Gly Ala Pro Glu Tyr Pro		
15	20	25
tct cag ctt gca gtt gaa atc gat cca gaa gcg att att gcg atc cag		208
Ser Gln Leu Ala Val Glu Ile Asp Pro Glu Ala Ile Ile Ala Ile Gln		
30	35	40

- 2 -

caa gat gca aac gcc gac cca cgt ctc ttt ttc cca ctg agc ggg ctt Gln Asp Ala Asn Ala Asp Pro Arg Leu Phe Phe Pro Leu Ser Gly Leu 45 50 55	256
gtc tcc gcc aaa ctt gcc aaa gtc ttt caa ccc aac ata tac cca acc Val Ser Ala Lys Leu Ala Lys Val Phe Gln Pro Asn Ile Tyr Pro Thr 60 65 70	304
cct cct agt ccc cag aca act tac cac ttt cac ctc cat cct cat ccc Pro Pro Ser Pro Gln Thr Thr Tyr His Phe His Leu His Pro His Pro 75 80 85 90	352
cat tat ccg cat cct cag cca agt tat cct cat cct caa ccc cat cat His Tyr Pro His Pro Gln Pro Ser Tyr Pro His Pro Gln Pro His His 95 100 105	400
cct cat cct cat tat cat cct cat cct cat ccc cat cat cct cat Pro His Pro His Pro Tyr His Pro His Pro His Pro His His Pro His 110 115 120	448
cct cat ccc cat caa cat cct cat cgt cat ccc gac cat cat ccc cac Pro His Pro His Gln His Pro His Arg His Pro Asp His His Pro His 125 130 135	496
cat cat cct cac cat cat cat gaa cat aat gtt cat gtg cct caa His His Pro His His His His Glu His Asn Val His Val Pro Gln 140 145 150	544
cat cag cac gct caa cac aac ggc cac cag aac aac ggt ggc cca gct His Gln His Ala Gln His Asn Gly His Gln Asn Asn Gly Gly Pro Ala 155 160 165 170	592
cat tat cac cat gac tac cat ttt gcg cat cct cat caa gag aac cag His Tyr His Asp Tyr His Phe Ala His Pro His Gln Glu Asn Gln 175 180 185	640
cat cac cgc gag gaa gag cag ctt acc gac atc aac taa gctattggtc His His Arg Glu Glu Gln Leu Thr Asp Ile Asn 190 195	689
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<212> PRT

<213> Eimeria tenella

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Ile Asp Pro Glu Ala Ile Ile Ala Ile Gln Gln Asp Ala Asn Ala Asp
35 40 45

Pro Arg Leu Phe Phe Pro Leu Ser Gly Leu Val Ser Ala Lys Leu Ala
50 55 60

Lys Val Phe Gln Pro Asn Ile Tyr Pro Thr Pro Pro Ser Pro Gln Thr.
65 70 75 80

Thr Tyr His Phe His Leu His Pro His Pro His Tyr Pro His Pro Gln
85 90 95

Pro Ser Tyr Pro His Pro Gln Pro His His Pro His Pro His Pro Tyr
100 105 110

His Pro His Pro His Pro His His Pro His Pro His Pro His Gln His
115 120 125

Pro His Arg His Pro Asp His His Pro His His His Pro His His His
130 135 140

His His Glu His Asn Val His Val Pro Gln His Gln His Ala Gln His
145 150 155 160

Asn Gly His Gln Asn Asn Gly Gly Pro Ala His Tyr His His Asp Tyr
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His Phe Ala His Pro His Gln Glu Asn Gln His His Arg Glu Glu Glu
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Gln Leu Thr Asp Ile Asn
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<213> *Eimeria tenella*

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atccagcaag atgcaaacgc cgaccacgt ctcttttcc cactgagcgg gtttgtctcc	180
gccaaacttg ccaaagtctt tcaacccaac atataaccaa cccctccatg tccccagaca	240
acttaccact ttacacctcca tcctcatccc cattatccgc atcctcagcc aagttatcct	300
catectcaac cccatcatcc tcatcctcat ctttatcatc ctcatcctca tccccatcat	360
cctcatacctc atccccatca acatcctcat cgtcatcccg accatcatcc ccaccatcat	420
cctcaccatc atcatcatga acataatgtt catgtgcctc aacatcagca cgctcaacac	480
aacggccacc agaacaacgg tggcccagct cattatcacc atgactacca ttttgcgcac	540
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<223> Primer A17-112-lo

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<210> 7

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<211> 25

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<400> 8

ctgtgagaag aaccgggtgc tcttc

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<210> 9

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<223> Primer EtACTIN-1o

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<211> 16

<212> DNA

<213> Artificial Sequence

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<223> Primer M13 forward

<400> 22

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<223> Primer T7 promoter

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<210> 24

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<212> DNA

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<223> Primer BGH reverse

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tagaaggcac agtcgagg

18